



Towards diagnosis–treatment combinations

Hans Helder, Wageningen University (NL)



- ▶ Diagnosis–treatment combinations (DTC)
 - ▶ A relatively new concept in healthcare
 - ▶ in The Netherlands
- ▶ (On January 1, 2005, it was introduced as a new financing system for hospitals).

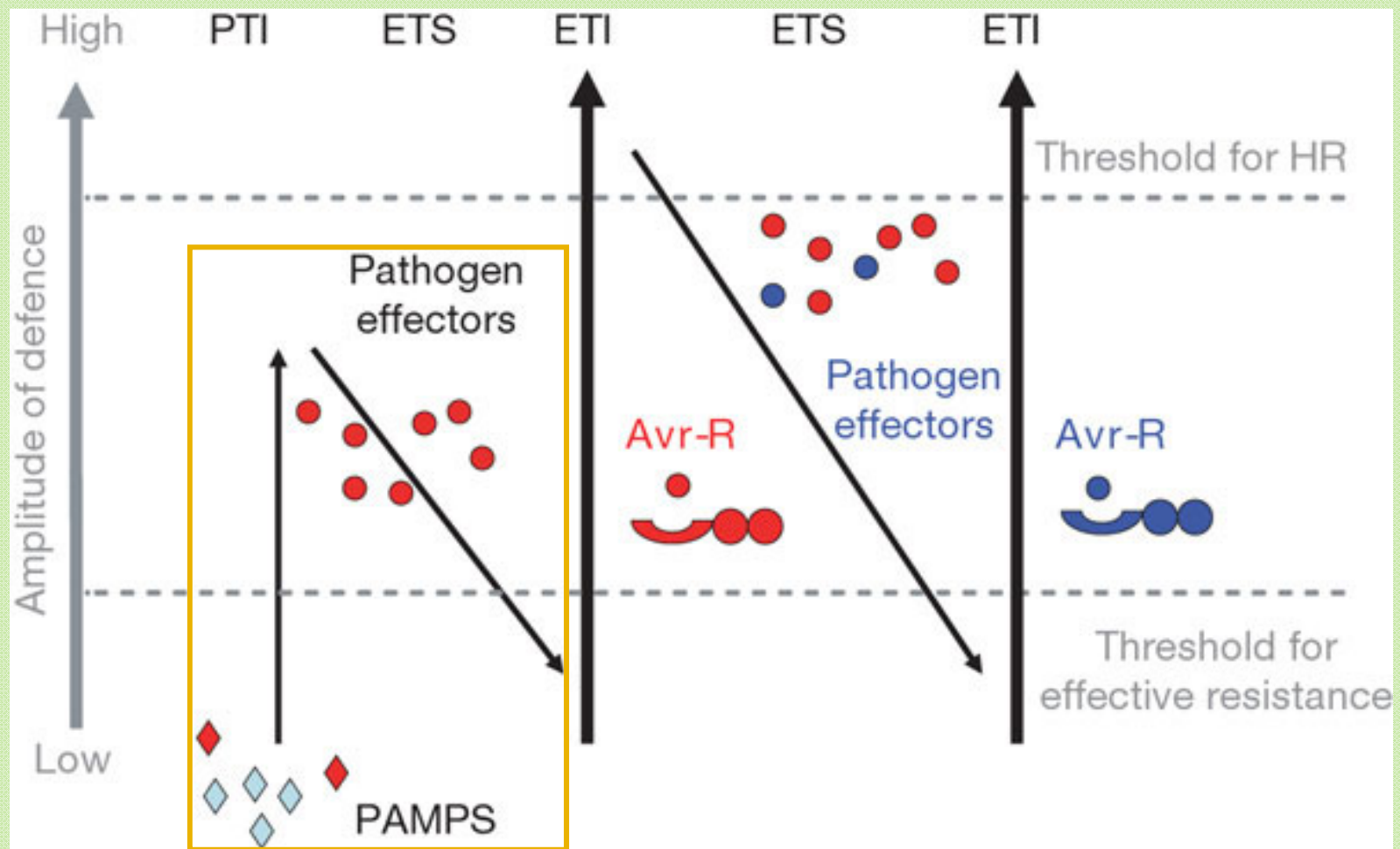


- ▶ A DTC includes all the activities and actions performed by the hospital and medical specialist in response to the patient's need for care, from the first consultation or examination to the final checkup.
- ▶ A DTC gives better insight into treatments performed by hospitals and the cost of such treatments, so that healthcare insurers know exactly what they are paying for and can compare hospitals and specializations.



Diagnosis – treatment combinations in plant pathology

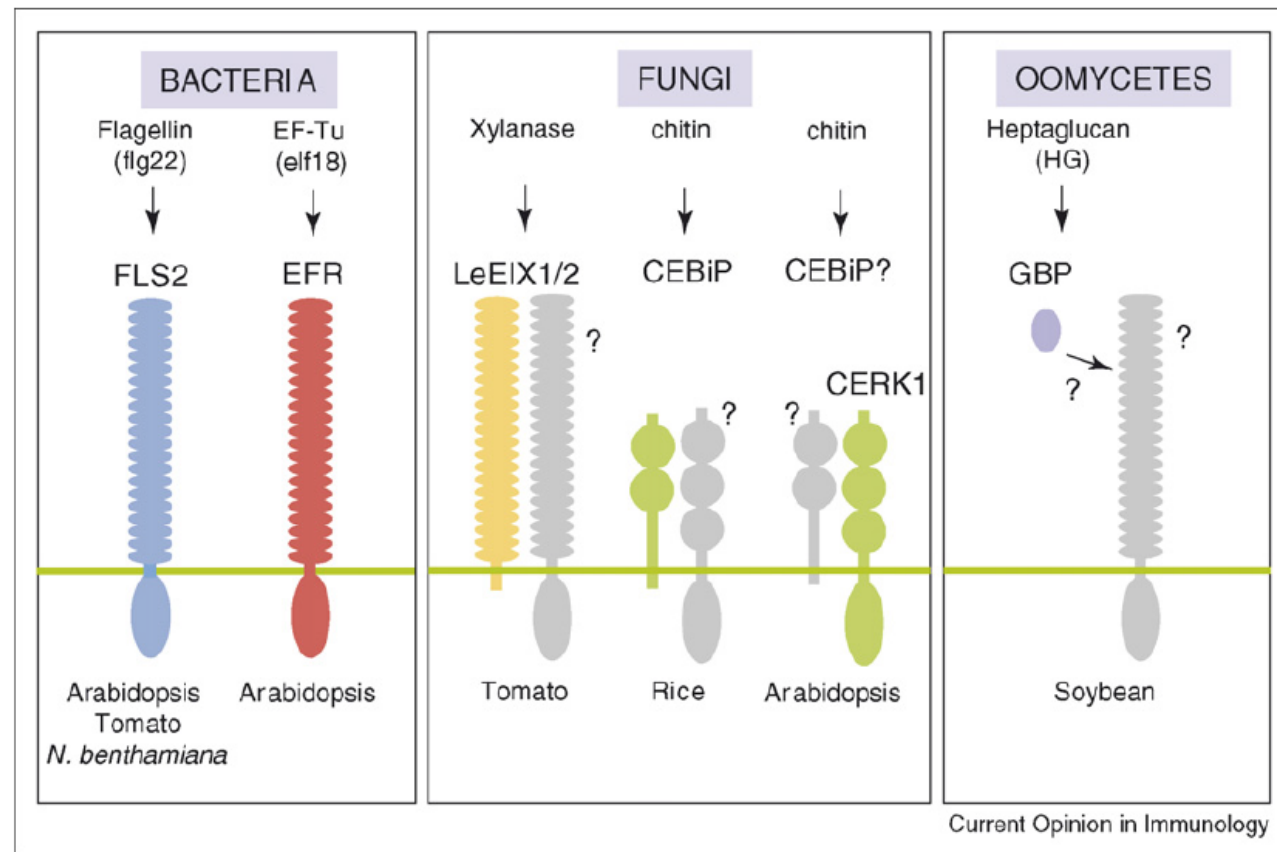
Diagnosis: Pathogens		Treatment: Handles in IPM
virus		Culture measures (<i>e.g.</i> (non-) tillage, crop rotation, etc.)
bacteria		PAMP-triggered immunity (PTI) Basal resistance
phytoplasmas		Effector-triggered immunity (ETI) Host plant resistance
oomycetes		Management general and specific (soil) suppressiveness
fungi		Biological control
nematodes		Pheromones
insects		<i>Agrochemicals</i>



Jones and Dangl (2006)

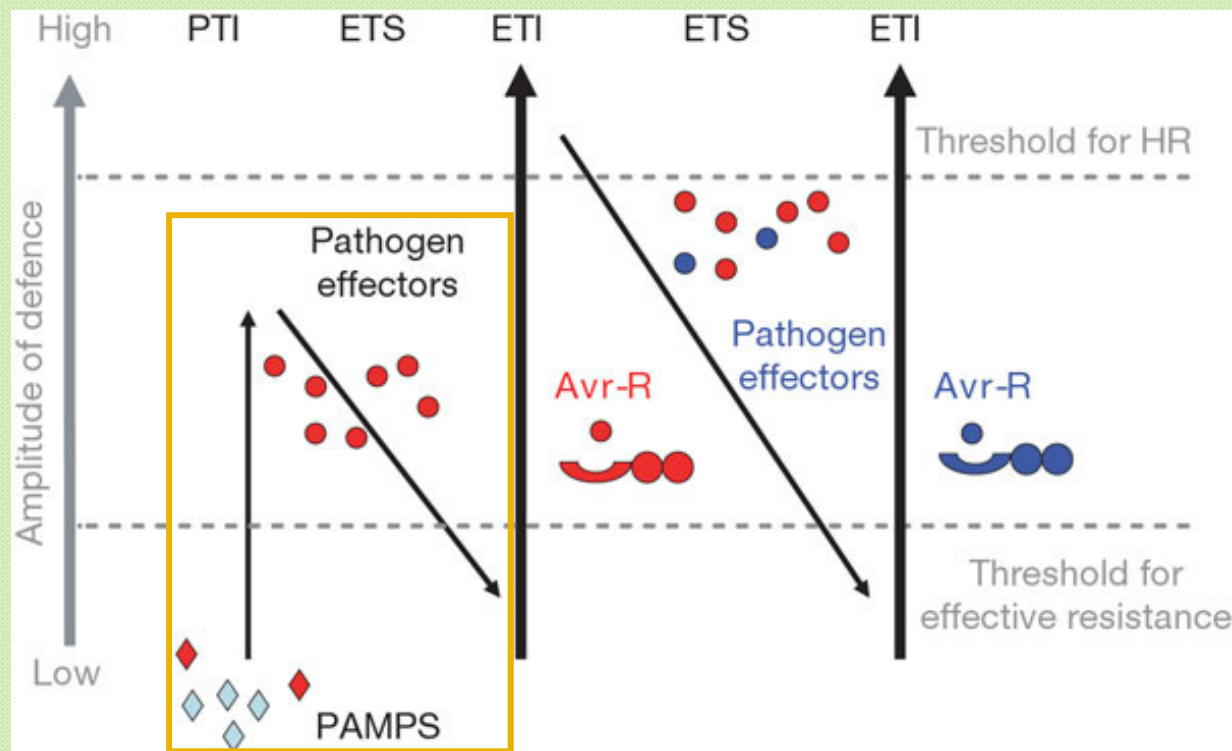
1. Plants detect microbial/pathogen-associated molecular patterns (MAMPs/PAMPs, red diamonds) via PRRs to trigger PAMP-triggered immunity (PTI) that can halt further colonization.

PRRs: transmembrane pattern recognition receptors



Cyril Zipfel (2008)

Plant PRRs. Bacterial flagellin (flg22) and EF-Tu (elf18) are recognized by the LRR-RLKs FLS2 and EFR, respectively. Orthologues of FLS2 have been cloned and characterized in tomato and *N. benthamiana*. In tomato, xylanase is recognised by the RLPs LeEIX1 and LeEIX2. LRR: leucine-rich repeat (leucine: hydrophobic amino acid); RLK: receptor-like protein kinase

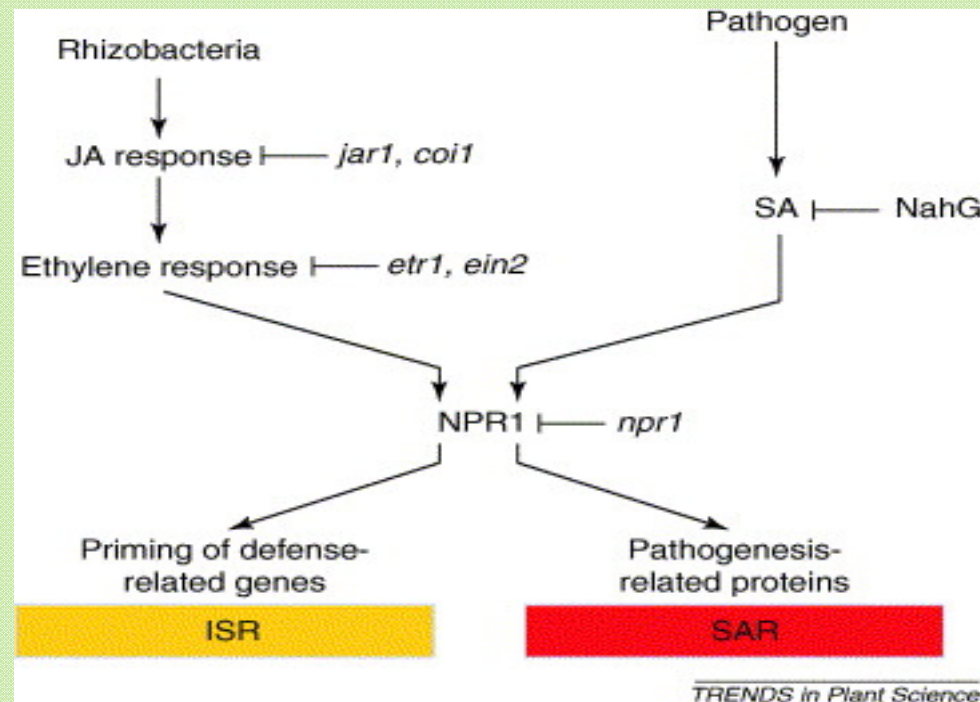


Jones and Dangl (2006)

PAMP-triggered immunity (PTI) –formerly named
basal or non-cultivar-specific resistance
 (to be compared with animal innate immunity).

= Ancient & conserved first layer of defense, it is based on the perception of conserved microbial structures by PRRs and it is effective against a broad spectrum of invading microorganisms

How to measure PAMP-triggered immunity (PTI)? NPR1 as an example



Signal transduction pathways of SAR and ISR.

SAR – Systemic acquired resistance (induced by an avirulent pathogen or upon restricted infection by a virulent pathogen)

ISR – Induced systemic resistance (induced by non-pathogenic rhizobacteria)

NPR1 – non-expressor of PR genes

A functional NPR1 protein is required for ISR and SAR – marker for basal resistance

From: L.C. van Loon, Bart P.J., Geraats, Huub J.M., Linthorst (2006)
Trends in Plant Science 11 (4) 184 – 191

OK, but are these *Arabidopsis* findings relevant for crops?

RESEARCH PAPER

***Malus hupehensis* NPR1 induces pathogenesis-related protein gene expression in transgenic tobacco**

J.-Y. Zhang^{1,2}, Y.-S. Qiao¹, D. Lv¹, Z.-H. Gao¹, S.-C. Qu¹ & Z. Zhang¹

¹ College of Horticulture, Nanjing Agricultural University, Nanjing, China

² Institute of Botany, Jiangsu Province and the Chinese Academy of Sciences, Nanjing, China

Research article

Open Access

Characterization of *Vitis vinifera* NPR1 homologs involved in the regulation of Pathogenesis-Related gene expression

Gaëlle Le Henanff¹, Thierry Heitz², Pere Mestre³, Jérôme Mutterer², Bernard Walter¹ and Julie Chong*¹

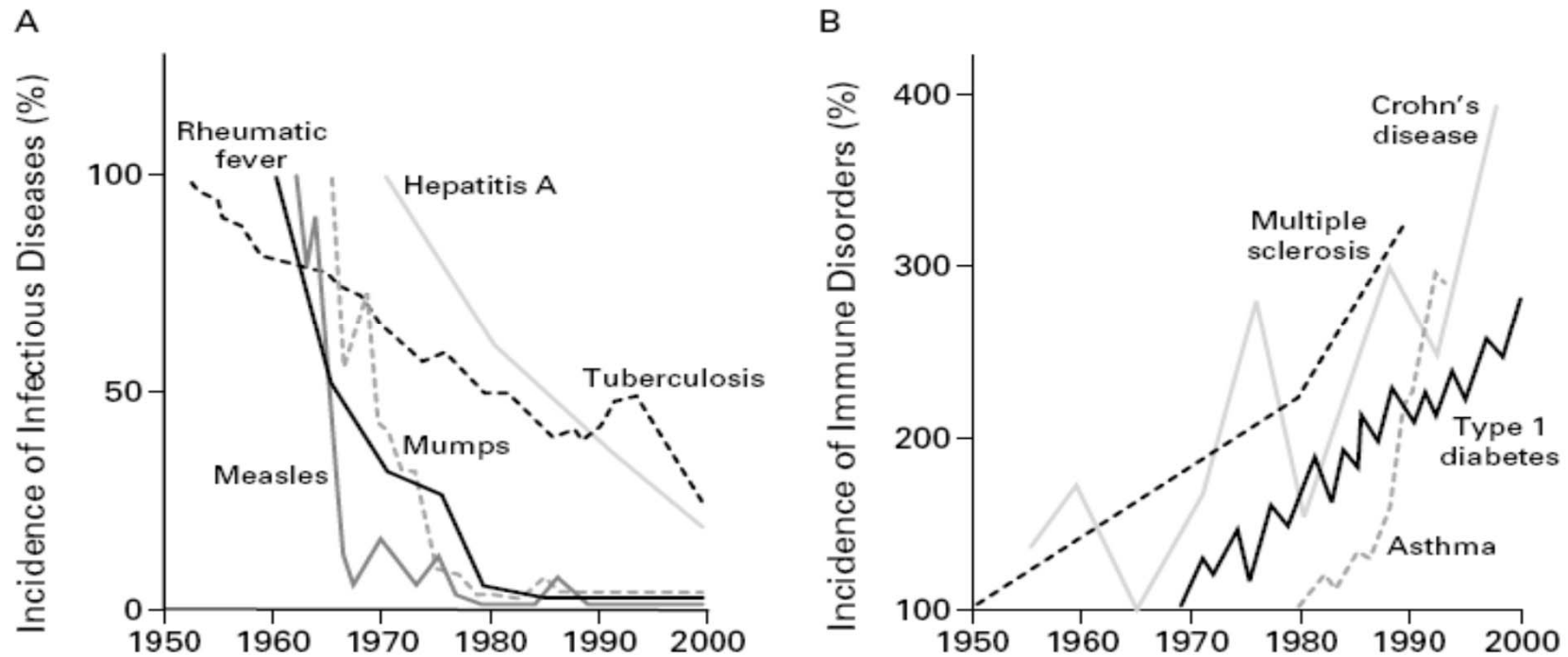
Functional analysis of the *Theobroma cacao* NPR1 gene in *arabidopsis*

Zi Shi¹, Siela N Maximova², Yi Liu¹, Joseph Verica², Mark J Guiltinan^{1,2*}

Etc., etc.

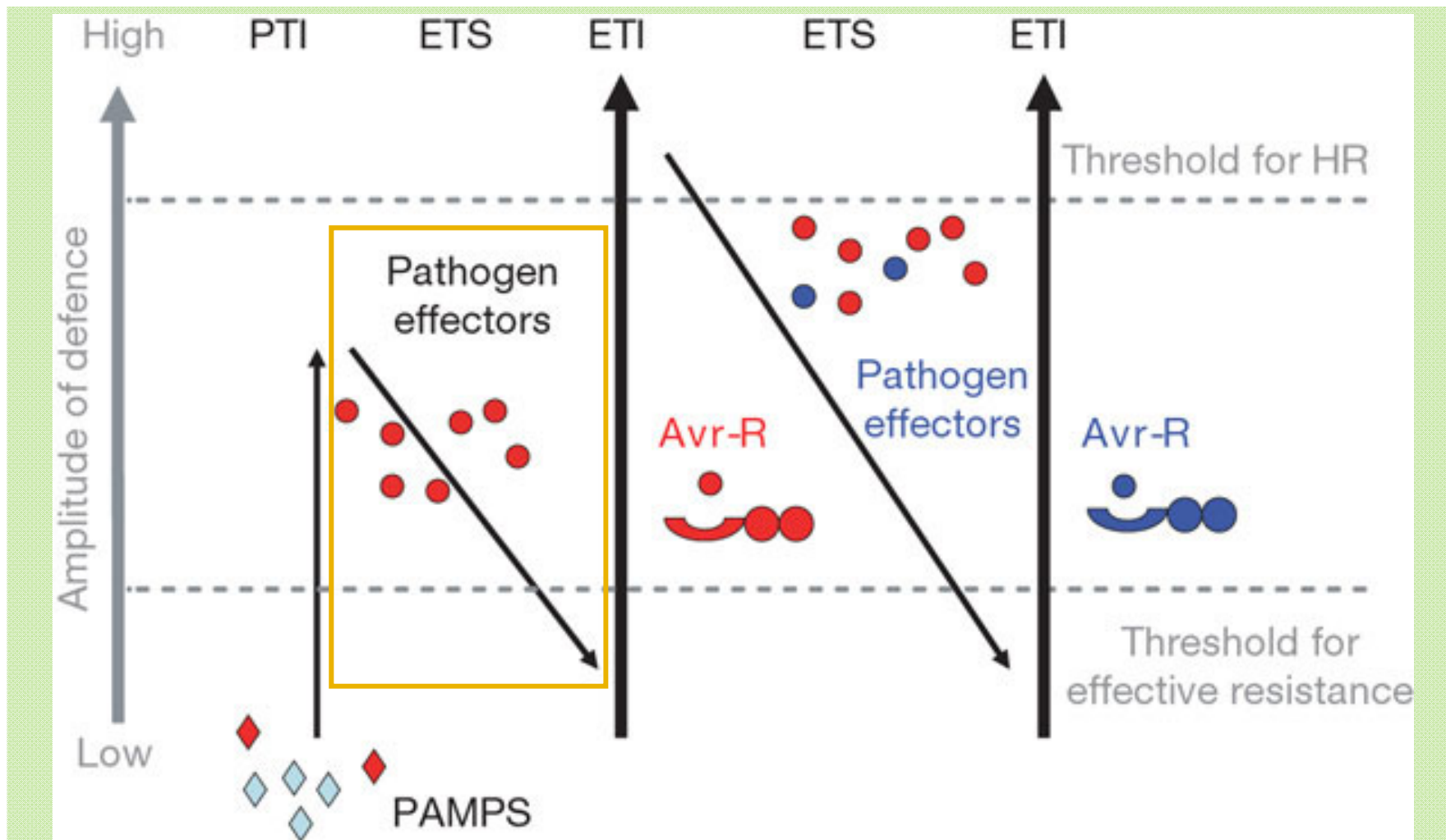


Inverse relation infectious diseases and immune disorders: hygiene hypothesis

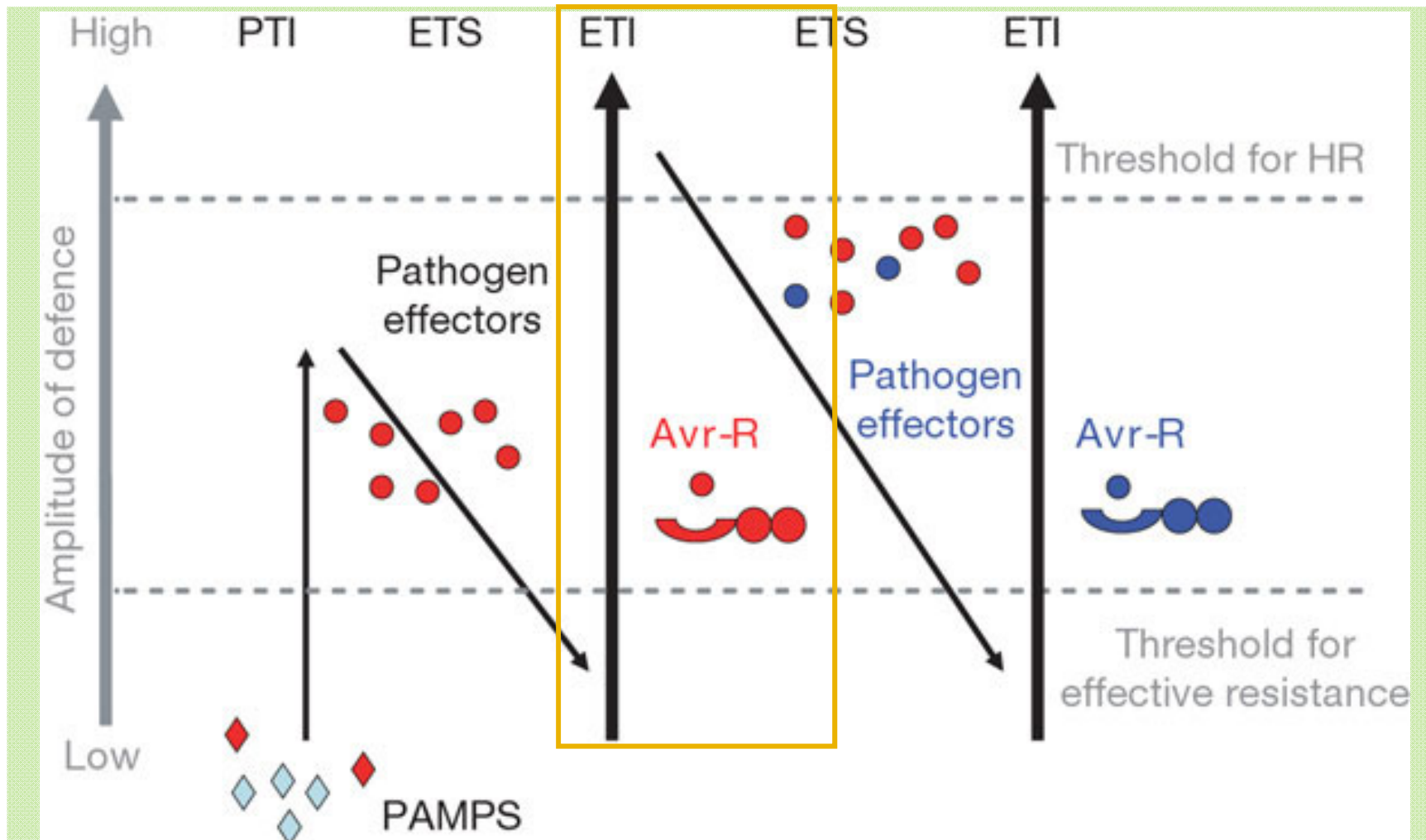


N Engl J Med 2002

The developing immune system must receive stimuli (from infectious agents, symbiotic bacteria, or parasites) in order to adequately develop regulatory T cells

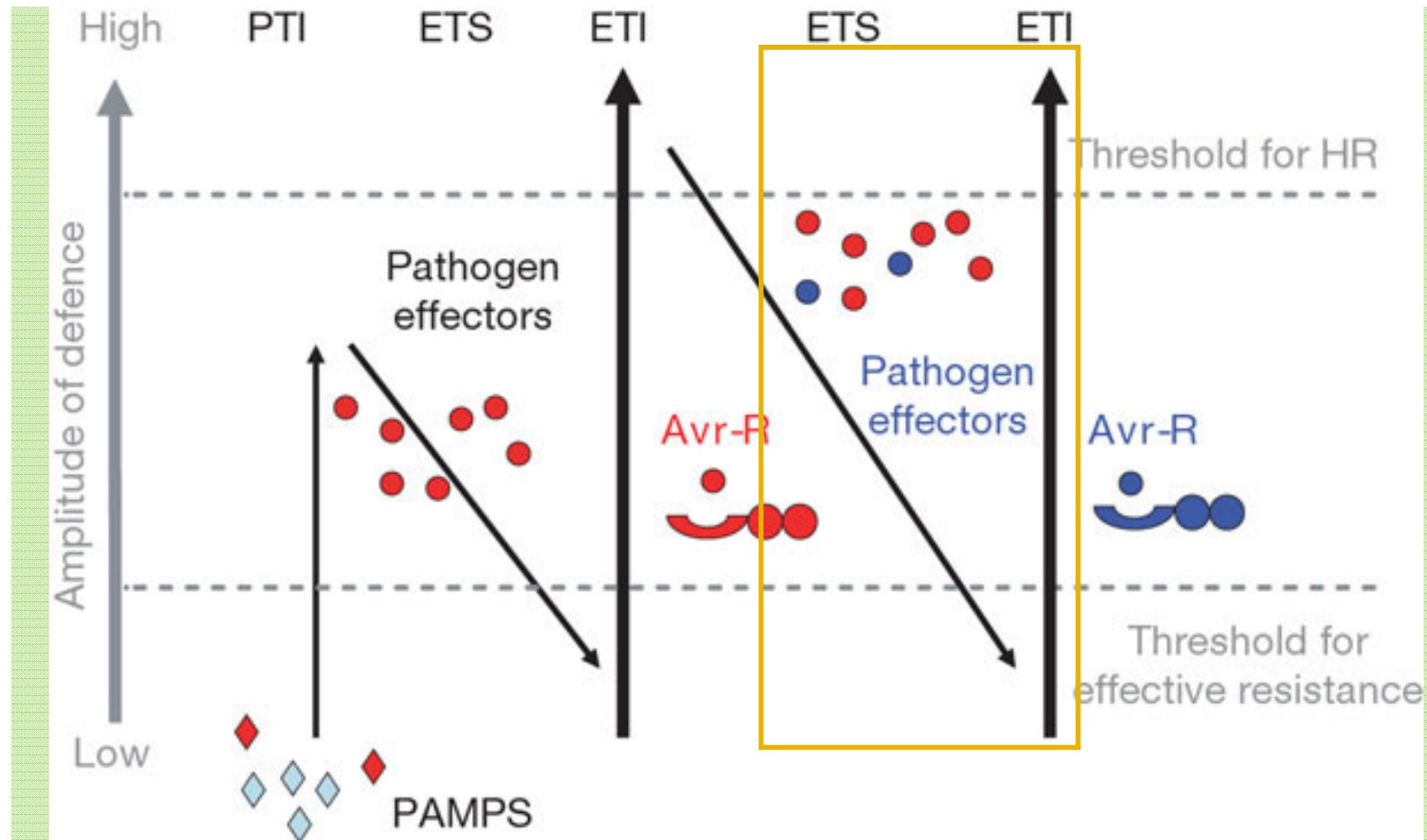


2. Successful pathogens deliver effectors that interfere with PTI, or otherwise enable pathogen nutrition and dispersal, resulting in effector-triggered susceptibility (ETS).



3. A given effector is 'specifically recognized' by one of the NB-LRR proteins, resulting in effector-triggered immunity (ETI).

ETI is an accelerated and amplified PTI response



Jones and Dangl (2006)

4. Natural selection drives pathogens to avoid ETI either by shedding or diversifying the recognized effector gene, or by acquiring additional effectors that suppress ETI. Natural selection results in new *R* specificities so that ETI can be triggered again.

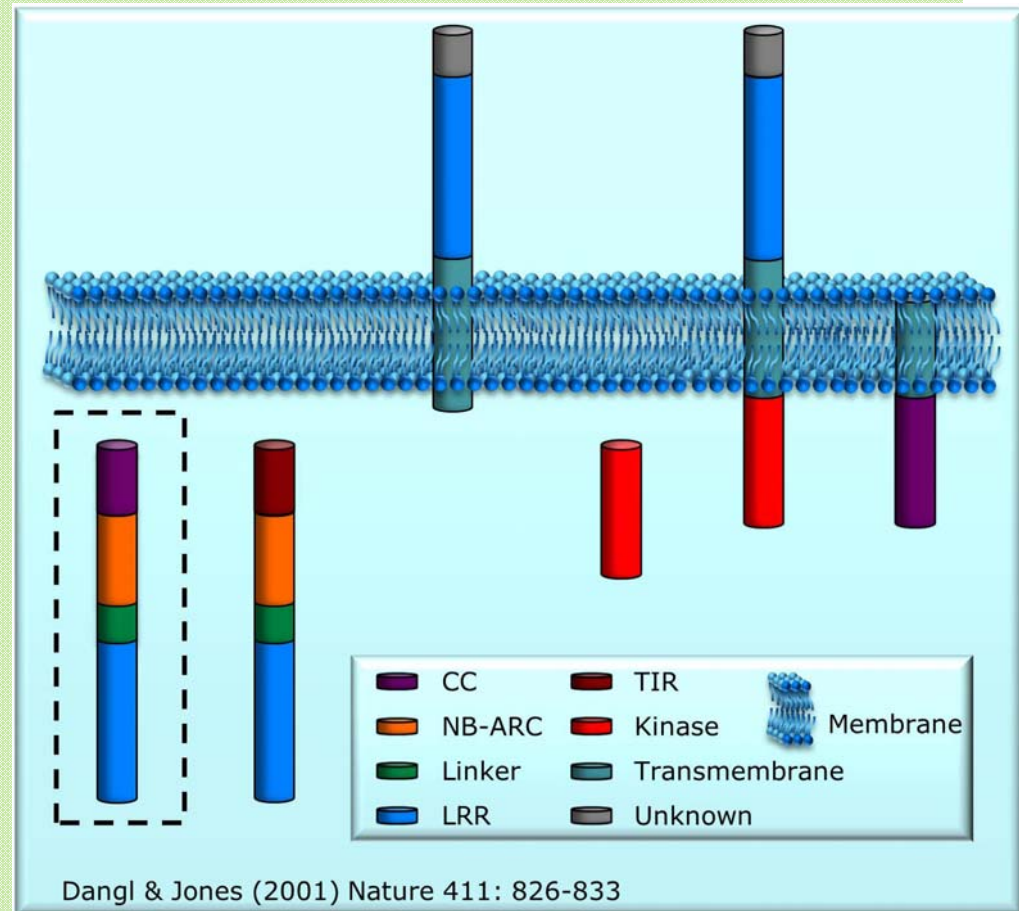
Effector-triggered immunity (ETI)

Resistance proteins

- highly **specific**
- strong response
- divided in classes based on their **domains**

NB-ARC-LRR

- the most common R proteins
- ~ **800** in **potato** genome
- N-terminal **CC** or **TIR**



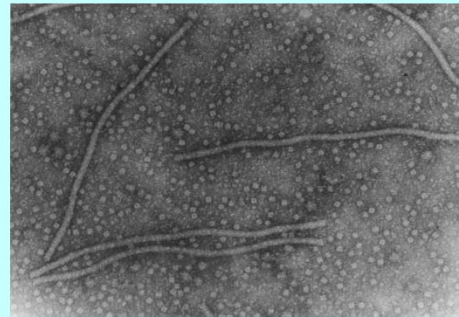
ETI, Example 1: virus and nematode

Two highly similar R proteins recognize two very different pathogens

Rx1

Extreme resistance to
Potato Virus X

Elicitor: PVX Coat protein



Virus particle: l = 515 nm
□ = 13 nm



PVX symptoms on
N. benthamiana

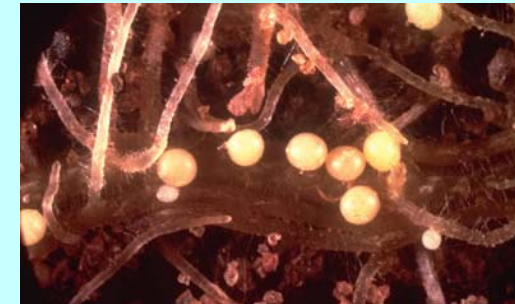
Gpa2

Slow response to
G. pallida D383

Elicitor: secreted
RanBP-like protein



Length ~500 µm



Cysts on roots
www.eppo.org

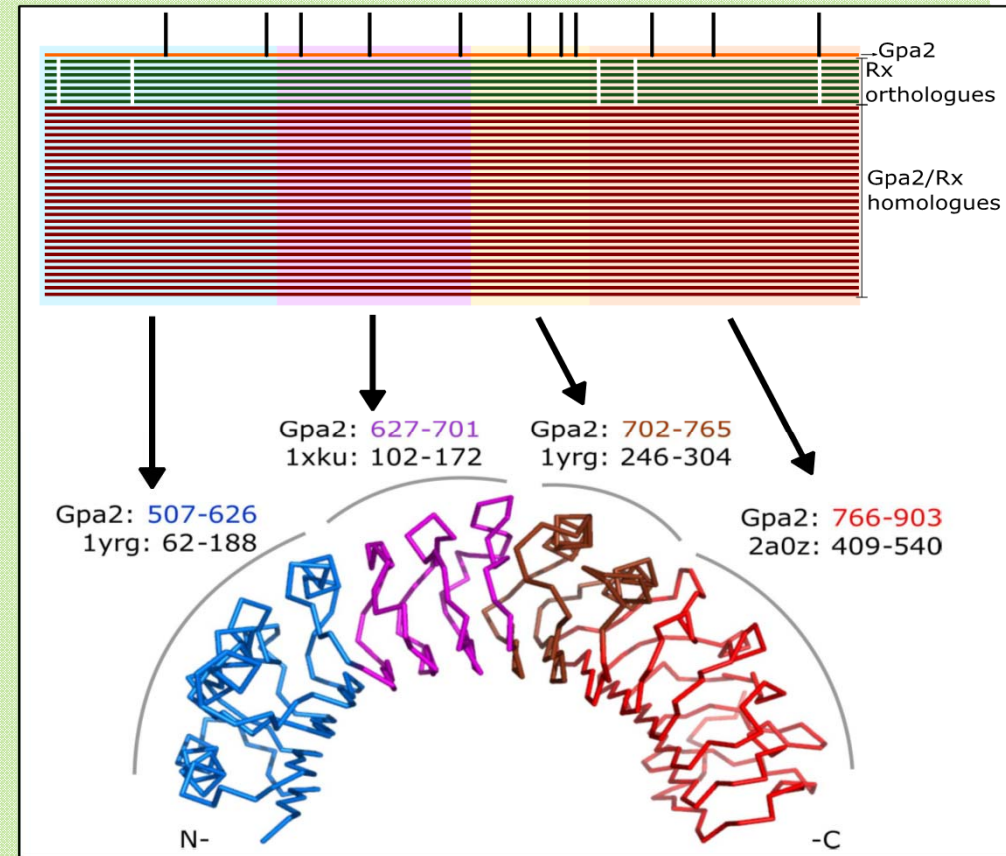
ETI, Example 1: virus and nematode

Full length dataset used to calculate 3D model of Gpa2/Rx LRR

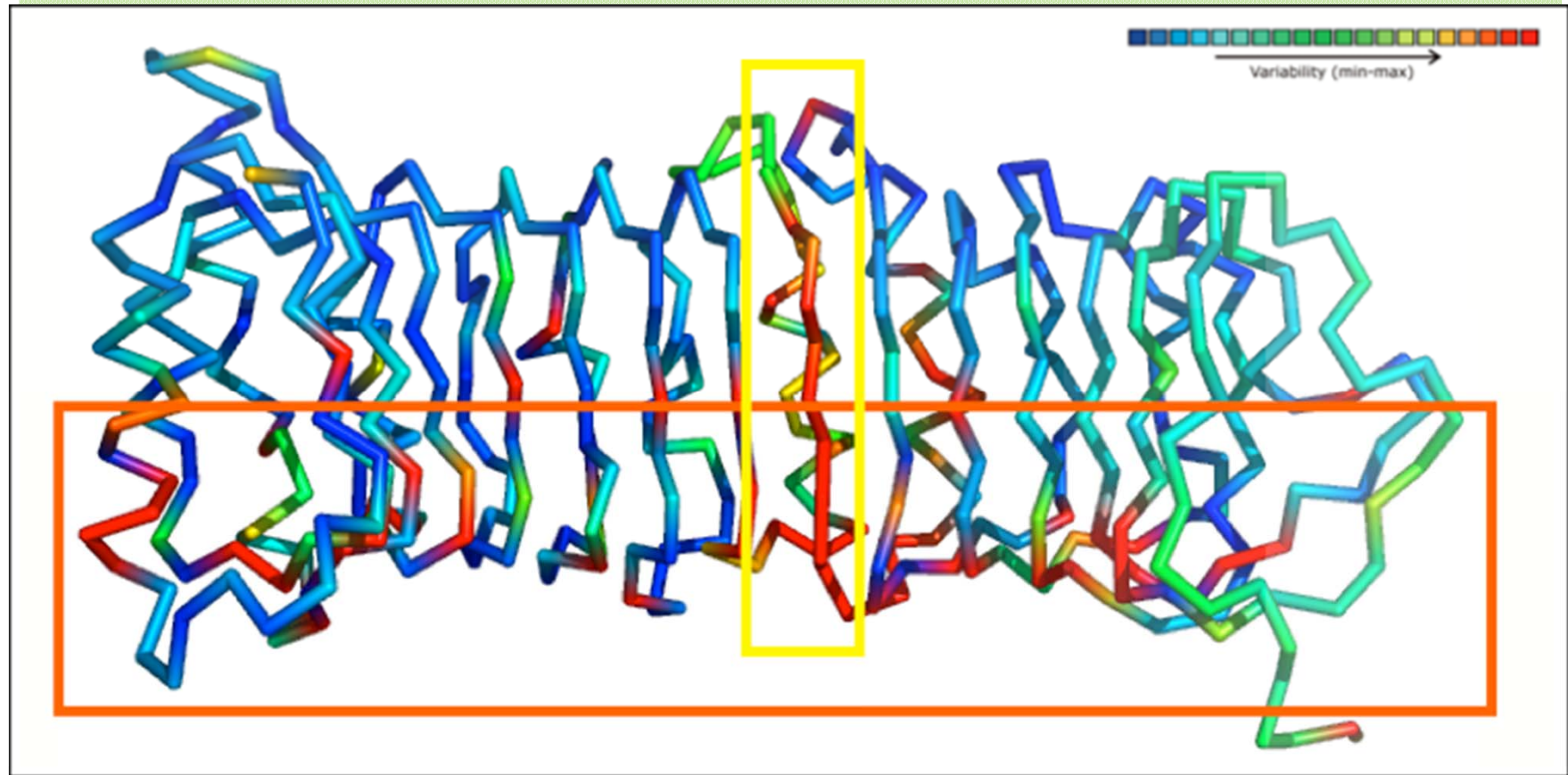
Gpa2 LRR is modeled after multiple templates

- Too irregular for single template
- Best structural template chosen per LRR segment

- RanGAP rna1p (1yrg),
- TLR3 (2a0z),
- Decorin (1xku)



ETI, Example 1: virus and nematode
Diversity of Gpa2/Rx homologs



How to build a pathogen detector: structural basis of NB-LRR function

Frank LW Takken^{1,3} and Aska Govere^{2,3}

Available online at www.sciencedirect.com

SciVerse ScienceDirect

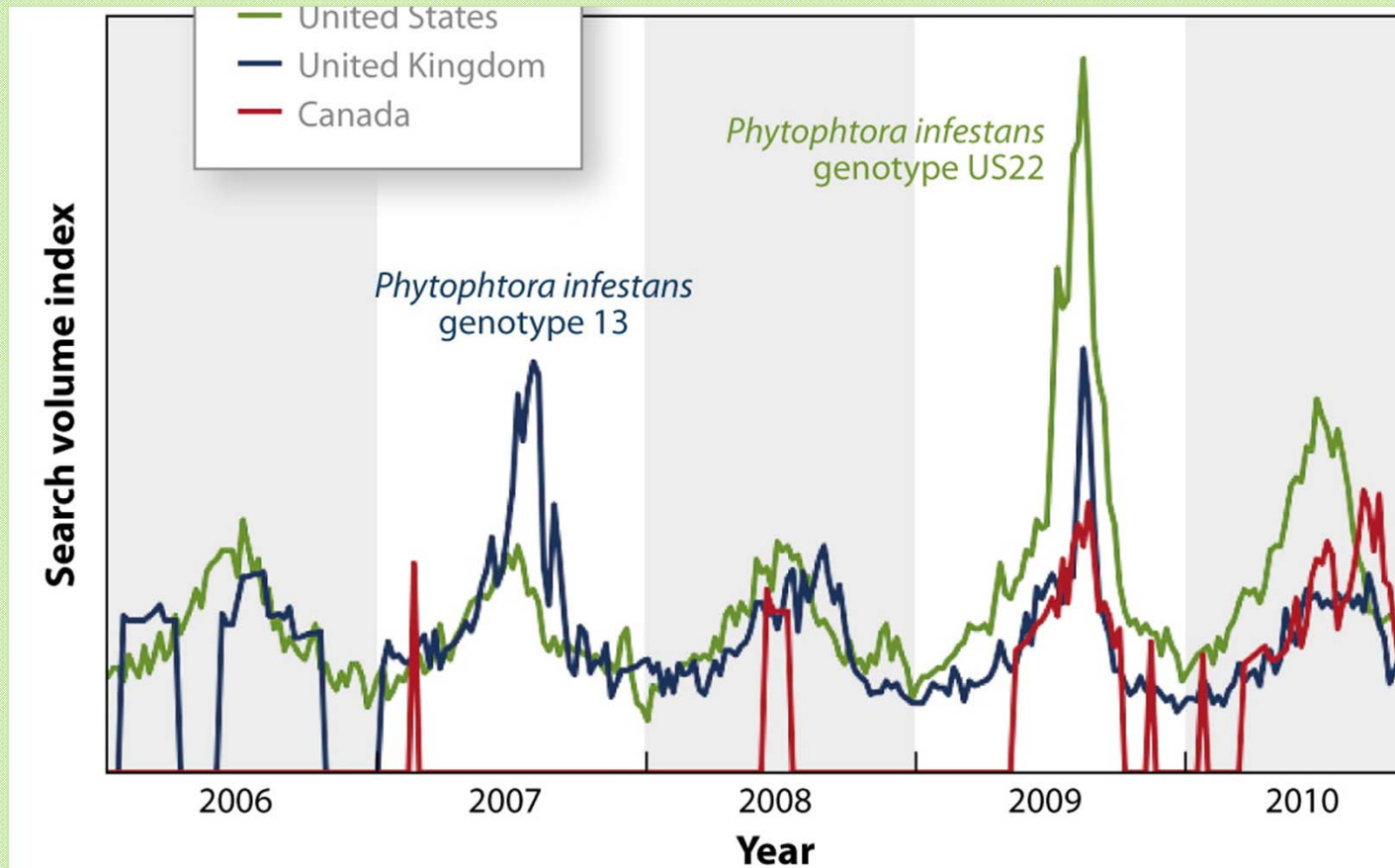
Current Opinion in
Plant Biology



ETI, Example 2: oomycete



ETI – Example 2 - oomycete



 Vleeshouwers VGAA, et al. 2011.

Numerous R genes available in potato that confer resistance against late blight

Table 1 *Solanum* R genes that confer resistance to *Phytophthora infestans*

R gene	RGH ^a	<i>Solanum</i> species ^b	Origin ^c	Chr ^d	Reference
R1 family	None				
<i>R1</i>		<i>demissum</i>	Mexico	V	(3, 86)
R2 family	None				
<i>R2</i>		<i>demissum</i>	Mexico	IV	(8, 78, 82)
<i>Rpi-blb3</i>		<i>bulbocastanum</i>	Mexico	IV	(78, 95)
<i>Rpi-abpt</i>		Unknown ^e	Mexico	IV	(49, 78, 97)
<i>R2-like</i>		<i>edinense</i>	Mexico	IV	(17, 78, 96)
<i>Rpi-edn1.1</i>		<i>edinense</i>	Mexico	IV	(17)
<i>Rpi-snk1.1</i>		<i>schenckii</i>	Mexico	IV	(17)
<i>Rpi-snk1.2</i>		<i>schenckii</i>	Mexico	IV	(17)
<i>Rpi-hjt1.1</i>		<i>hjertingii</i>	Mexico	IV	(17)
<i>Rpi-hjt1.2</i>		<i>hjertingii</i>	Mexico	IV	(17)
<i>Rpi-hjt1.3</i>		<i>hjertingii</i>	Mexico	IV	(17)
<i>Rpi-mcd1</i>		<i>microdontum</i>	Argentina	IV	(77)
R3a family	I2 (94)				
<i>R3a</i>		<i>demissum</i>	Mexico	XI	(55, 56)
<i>Rpi-sto2</i>		<i>stoloniferum</i>	Mexico	XI	(17)
R4 family	Unknown ^f				
<i>R4</i>		<i>demissum</i>	Mexico	XI	(136, 142)
<i>Rpi-blb1</i> family	None				
<i>Rpi-blb1, RB</i>		<i>bulbocastanum</i>	Mexico	VIII	(123, 134)
<i>Rpi-sto1</i>		<i>stoloniferum</i>	Mexico	VIII	(145)
<i>Rpi-pta1</i>		<i>stoloniferum</i> ^g	Mexico	VIII	(145)
<i>Rpi-blb2</i> family	Mi (114)				
<i>Rpi-blb2</i>		<i>bulbocastanum</i>	Mexico	VI	(135)
<i>Rpi-vnt1</i> family	Tm2 ² (72)				
<i>Rpi-vnt1.1</i>		<i>venturii</i>	Argentina	IX	(100)
<i>Rpi-vnt1.2</i>		<i>venturii</i>	Argentina	IX	(30)
<i>Rpi-vnt1.3</i>		<i>venturii</i>	Argentina	IX	(100)

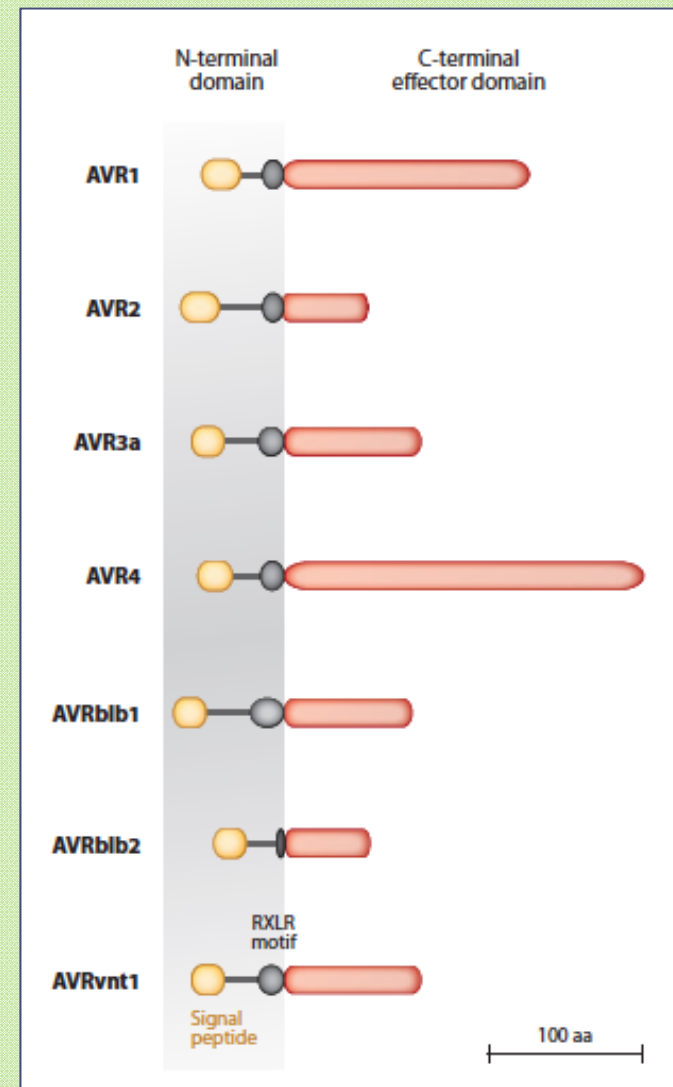
Vleeshouwers et al. 2011



Features of characterized *Phytophthora Avr* gene products

C-terminal domain carries the effector biochemical activity

Vleeshouwers et al. 2011,
Ann Rev Phytopathol 49: 507–



IPM compatible, environmentally sound solutions such as

- PTI
- ETI
- biocontrol
- pheromone-based control measures

are feasible but almost inevitably **knowledge intensive**

Pathogens should be diagnosed early

**in most cases at least at species level, and preferably (for ETI)
at effector level ('host race', 'pathovar', 'pathotype' etc.)**

in a high throughput, affordable and fast manner

DNA-based diagnostics seem most appropriate to reach these goals

Requirements for building large DNA frameworks:

1. Thorough classical (phytopathological)
taxonomical knowledge

(scarce)

2. Molecular expertise

(less scarce)

3. Perseverance & some recourses

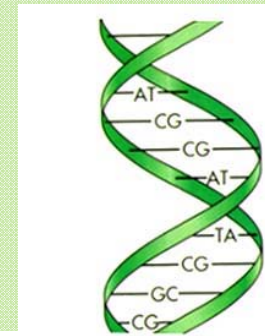
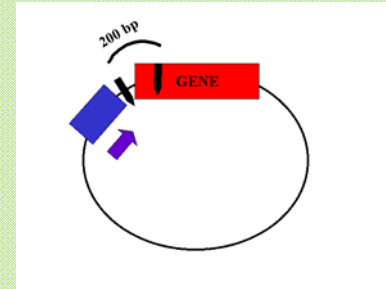
(increasingly scarce)

From morphology to DNA-based diagnostics



Parasitic and non-parasitic
nematodes

≈ 2,500 taxa



Small Subunit Ribosomal DNA-Based Phylogenetic Analysis of Foliar Nematodes (*Aphelenchoides* spp.) and Their Quantitative Detection in Complex DNA Backgrounds

Katarzyna Rybarczyk-Mydlowska, Paul Mooyman, Hanny van Megen, Sven van den Elsen, Mariëtte Vervoort, Peter Veenhuizen, Joop van Doorn, Robert Dees, Gerrit Karssen, Jaap Bakker, and Johannes Helder

Phytopathology 2012

+ assays for cyst, root knot, lesion, stem and stubby root nematodes

Next major step: **effector-based pathogen diagnostics**

Maximizing the use basal and effector-based plant immunity
by match making

Akbar Karegar (Shiraz, Iran)
Wim Bert (Ghent, Belgium)
Gerrit Karssen (NRL)
Joop van Doorn (PPO Lisse)
Jet Vervoort
Hanny van Megen
Sven van den Elsen
Paul Mooyman
Kasia Rybarczyk-Mydłowska
Casper Quist
Erik Slootweg
Erin Bakker
Geert Smant
Aska Goverse
Jaap Bakker

Laboratorium of Nematology, Wageningen University